



## SEQUENCE LISTING

<110> Marasco, Wayne  
Marasco, Abner

<120> INTRABODY-MEDIATED CONTROL OF IMMUNE REACTIONS

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<140> 09/522,727

<141> 2000-03-10

<150> PCT/US98/19563

<151> 1998-09-18

<150> 60/059,339

<151> 1997-09-18

<160> 55

<170> PatentIn version 3.0

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&lt;212&gt; PRT

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Glu Gly Lys Ser Ser Gly Ser Gly Ser Glu Ser Lys Val Asp  
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&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 6

Gly Ser Thr Ser Gly Ser Gly Lys Ser Ser Glu Gly Lys Gly  
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&lt;211&gt; 15

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 7

Lys Glu Ser Gly Ser Val Ser Ser Glu Gln Leu Ala Gln Phe Arg  
 1 5 10 15

&lt;210&gt; 8

&lt;211&gt; 16

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 8

Glu Ser Gly Ser Val Ser Ser Glu Glu Leu Ala Phe Arg Ser Leu Asp  
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&lt;210&gt; 9

&lt;211&gt; 35

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 9

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&lt;212&gt; DNA

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Asp Glu Glu Leu  
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Gln Glu Asp Leu  
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Arg Asp Glu Leu  
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Gly Gln Thr Val Thr Thr Pro Leu  
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<210> 32

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Gly Gln Glu Leu Ser Gln His Glu  
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Gly Gln Thr Leu Thr Thr Pro Leu  
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Gly Gln Ile Phe Ser Arg Ser Ala

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EI  
 Conf

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Gly Cys Thr Leu Ser Ala Glu Glu  
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Gly Ala Gln Leu Ser Arg Asn Thr  
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Gly Asn Ala Ala Ala Ala Lys Lys  
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Gly Asn Glu Ala Ser Tyr Pro Leu  
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Gly Ser Ser Lys Ser Lys Pro Lys  
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&lt;213&gt; Homo sapiens

&lt;400&gt; 49

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&lt;210&gt; 50

&lt;211&gt; 25

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 50

tgmggagacg gtgaccrwgg tccct

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&lt;210&gt; 51

&lt;211&gt; 837

&lt;212&gt; DNA

&lt;213&gt; human

&lt;220&gt;

&lt;221&gt; CDS

UNSURE

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(169)(505)

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5 10 15

gtc ctg tcc cag gtg caa ctg cag cag tca ggg gct gag ctg gca aga 96

Val Leu Ser Gln Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Ala Arg

20 25 30

cct ggg gct tca gtg aag ttg tcc tgc aag gct tct ggc tac acc ttt 144

Pro Gly Ala Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe

35 40 45

act agt cac tgg atg cag tgg gtg aga cag agg cct gga cag ggt ctg 192

Thr Ser His Trp Met Gln Trp Val Arg Gln Arg Pro Gly Gln Gly Leu

50 55 60

gaa tgg att ggg act att tat cct gga gat ggt gat act agg tac act 240

Glu Trp Ile Gly Thr Ile Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr

65 70 75 80

cag aat ttc aag ggc aag gcc aca ttg act gca gat aag tcc tcc acc 288

Gln Asn Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Thr

85 90 95

aca gcc tac tta cac ctc agc agc ttg tca tct gaa gac tct gcg gtc 336

Thr Ala Tyr Leu His Leu Ser Ser Leu Ser Ser Glu Asp Ser Ala Val

100 105 110

tat tat tgt gca aga gat gag att act acg gtt gta ccc cgg ggg ttt 384

Tyr Tyr Cys Ala Arg Asp Glu Ile Thr Thr Val Val Pro Arg Gly Phe

115 120 125

gct tac tgg ggc caa ggg acc tcg gtc acc gtc tcc tca ggt ggc ggt 432

Ala Tyr Trp Gly Gln Gly Thr Ser Val Thr Val Ser Ser Gly Gly Gly

130 135 140

ggc tcg ggc ggt ggt ggc tcg ggt ggc ggc gga tct gag ctc gtg ctc 480

Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Glu Leu Val Leu

145 150 155 160

acc caa acc cca acc tcc ctg gct ncc tct ctg gga gac aga gtc acc 528

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Thr Gln Thr Pro Thr Ser Leu Ala Xaa Ser Leu Gly Asp Arg Val Thr  
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 Ile Ser Cys Arg Ala Ser Gln Asp Ile Ser Ser Tyr Leu Asn Trp Tyr  
 180 185 190  
 cag cag aaa cca gat gga act att aaa ctc ctg atc tac tac aca tca 624  
 Gln Gln Lys Pro Asp Gly Thr Ile Lys Leu Leu Ile Tyr Tyr Thr Ser  
 195 200 205  
 aga tta tat tca gga gtc cca cca agg ttc agt ggc agt ggg gct gga 672  
 Arg Leu Tyr Ser Gly Val Pro Pro Arg Phe Ser Gly Ser Gly Ala Gly  
 210 215 220  
 aca gat tat tct ctc acc atc agc aac ctg gag caa gaa gat ttt gcc 720  
 Thr Asp Tyr Ser Leu Thr Ile Ser Asn Leu Glu Gln Glu Asp Phe Ala  
 225 230 235 240  
 act tac ttt tgc caa cag ggt aat gtg att ccg tac acg ttc gga ggg 768  
 Thr Tyr Phe Cys Gln Gln Gly Asn Val Ile Pro Tyr Thr Phe Gly Gly  
 245 250 255  
 ggg acc aag ctg gaa atg aaa cgg gct gat gct gca cca act gta agc 816  
 Gly Thr Lys Leu Glu Met Lys Arg Ala Asp Ala Ala Pro Thr Val Ser  
 260 265 270  
 gaa aag gac gag ctg taa taa 837  
 Glu Lys Asp Glu Leu  
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<400> 52

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 Pro Gly Ala Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe  
 35 40 45  
 Thr Ser His Trp Met Gln Trp Val Arg Gln Arg Pro Gly Gln Gly Leu  
 50 55 60  
 Glu Trp Ile Gly Thr Ile Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr  
 65 70 75 80  
 Gln Asn Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Thr  
 85 90 95  
 Thr Ala Tyr Leu His Leu Ser Ser Leu Ser Ser Glu Asp Ser Ala Val

Dry  
 cont  
 El  
 cont

100 105 110  
 Tyr Tyr Cys Ala Arg Asp Glu Ile Thr Thr Val Val Pro Arg Gly Phe  
 115 120 125  
 Ala Tyr Trp Gly Gln Gly Thr Ser Val Thr Val Ser Ser Gly Gly Gly  
 130 135 140  
 Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Glu Leu Val Leu  
 145 150 155 160  
 Thr Gln Thr Pro Thr Ser Leu Ala Xaa Ser Leu Gly Asp Arg Val Thr  
 165 170 175  
 Ile Ser Cys Arg Ala Ser Gln Asp Ile Ser Ser Tyr Leu Asn Trp Tyr  
 180 185 190  
 Gln Gln Lys Pro Asp Gly Thr Ile Lys Leu Leu Ile Tyr Tyr Thr Ser  
 195 200 205  
 Arg Leu Tyr Ser Gly Val Pro Pro Arg Phe Ser Gly Ser Gly Ala Gly  
 210 215 220  
 Thr Asp Tyr Ser Leu Thr Ile Ser Asn Leu Glu Gln Glu Asp Phe Ala  
 225 230 235 240  
 Thr Tyr Phe Cys Gln Gln Gly Asn Val Ile Pro Tyr Thr Phe Gly Gly  
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 Gly Thr Lys Leu Glu Met Lys Arg Ala Asp Ala Ala Pro Thr Val Ser  
 260 265 270  
 Glu Lys Asp Glu Leu  
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 <222> (1)...(837)  
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 1 5 10 15  
 gtc ctg tcc cag gtg caa ctg cag cag tct ggg gct gag ctg aca aga 96  
 Val Leu Ser Gln Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Thr Arg  
 20 25 30  
 cct ggg gct tca gtg aag ttg tcc tgc aag gct tct ggc tac acc ttt 144  
 Pro Gly Ala Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe  
 35 40 45  
 act agt cac tgg atg cag tgg gtg aga cag agg cct gga cag ggt ctg 192  
 Thr Ser His Trp Met Gln Trp Val Arg Gln Arg Pro Gly Gln Gly Leu  
 50 55 60

gaa tgg att ggg act att tat cct gga gat ggt gat act agg tac act 240  
 Glu Trp Ile Gly Thr Ile Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr  
 65 70 75 80  
 cag aat ttc aag ggc aag gcc aca ttg act gca gat aag tcc tcc acc 288  
 Gln Asn Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Thr  
 85 90 95  
 aca gcc tac tta cac ctg agc agc ttg tca tct gaa gac tct gcg gtc 336  
 Thr Ala Tyr Leu His Leu Ser Ser Leu Ser Ser Glu Asp Ser Ala Val  
 100 105 110  
 tat tat tgt gca aga gat gag att act acg gtt gta ccc cgg ggg ttt 384  
 Tyr Tyr Cys Ala Arg Asp Glu Ile Thr Thr Val Val Pro Arg Gly Phe  
 115 120 125  
 gct tac tgg ggc caa ggg acc ttg gtc acc gtc tcc tca ggt ggc ggt 432  
 Ala Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Gly Gly Gly  
 130 135 140  
 ggc tcg ggc ggt ggt ggg tcg ggt ggc ggc gga tct gag ctg gtg ctg 480  
 Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Glu Leu Val Leu  
 145 150 155 160  
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 Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Leu Gly Asp Thr Ile Thr  
 165 170 175  
 atc act tgc cat gcc agt cag aac att aat gtt tgg tta agt tgg tac 576  
 Ile Thr Cys His Ala Ser Gln Asn Ile Asn Val Trp Leu Ser Trp Tyr  
 180 185 190  
 cag cag aaa cca gga aat att cct caa cta ttg atc tat aag gct tcc 624  
 Gln Gln Lys Pro Gly Asn Ile Pro Gln Leu Leu Ile Tyr Lys Ala Ser  
 195 200 205  
 aac ttg cac aca ggc gtc cca tca agg ttt agt ggc cgt gga tct gga 672  
 Asn Leu His Thr Gly Val Pro Ser Arg Phe Ser Gly Arg Gly Ser Gly  
 210 215 220  
 aca ggt ttc aca tta acc atc agc agc ctg cag cct gaa gac att ggc 720  
 Thr Gly Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Gly  
 225 230 235 240  
 act tac tac tgt caa cag ggt caa agt tat cct ctg acg ttc ggt gga 768  
 Thr Tyr Tyr Cys Gln Gln Gly Gln Ser Tyr Pro Leu Thr Phe Gly Gly  
 245 250 255  
 ggc acc aag ctg gaa atc aaa cgg gct gat gct gca cca act gta agc 816  
 Gly Thr Lys Leu Glu Ile Lys Arg Ala Asp Ala Ala Pro Thr Val Ser  
 260 265 270  
 gaa aag gac gag ctg taa taa 837  
 Glu Lys Asp Glu Leu  
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Met Glu His Leu Trp Phe Phe Leu Leu Leu Val Ala Ala Pro Arg Trp  
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Pro Gly Ala Ser Val Leu Leu Ser Cys Leu Ala Ser Gly Tyr Thr Phe  
 35 40 45

Thr Ser His Trp Met Gln Trp Val Arg Gln Arg Pro Gly Gln Gly Leu  
 50 55 60

Glu Trp Ile Gly Thr Ile Tyr Pro Gly Asp Gly Asp Thr Arg Tyr Thr  
 65 70 75 80

Gln Asn Phe Leu Gly Leu Ala Thr Leu Thr Ala Asp Leu Ser Ser Thr  
 85 90 95

Thr Ala Tyr Leu His Leu Ser Ser Leu Ser Ser Glu Asp Ser Ala Val  
 100 105 110

Tyr Tyr Cys Ala Arg Asp Glu Ile Thr Thr Val Val Pro Arg Gly Phe  
 115 120 125

Ala Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Gly Gly Gly  
 130 135 140

Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Glu Leu Val Leu  
 145 150 155 160

Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Leu Gly Asp Thr Ile Thr  
 165 170 175

Ile Thr Cys His Ala Ser Gln Asn Ile Asn Val Trp Leu Ser Trp Tyr  
 180 185 190

Gln Gln Leu Pro Gly Asn Ile Pro Gln Leu Leu Ile Tyr Leu Ala Ser  
 195 200 205

Asn Leu His Thr Gly Val Pro Ser Arg Phe Ser Gly Arg Gly Ser Gly  
 210 215 220

Thr Gly Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro Glu Asp Ile Gly  
 225 230 235 240

Thr Tyr Tyr Cys Gln Gln Gly Gln Ser Tyr Pro Leu Thr Phe Gly Gly  
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 260 265 270

Glu Leu Asp Glu Leu  
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<400> 55

Arg Lys Lys Arg  
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